

**Evaluating the dietary micro-remain record in dental calculus and its application in deciphering hominin diets in Palaeolithic Eurasia** Power, R.C.F.

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## **Assessing use and suitability of scanning electron microscopy in the analysis of microremains in dental calculus**

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#### **Abstract**

Dental calculus is increasingly recognised as a major reservoir of dietary information. Palaeodietary studies using plant and animal microremains (e.g. phytoliths, pollen, sponge spicules, and starch grains) trapped in calculus have the potential to revise our knowledge of the dietary role of plants in past populations. The conventional methods used to isolate and identify these microremains rely on removing them from their microenvironment in the calculus, thus the microenvironment that traps and preserves microremains is not understood. By using scanning electron microscopy and energy-dispersive X-ray spectroscopy (SEM-EDX) on modern chimpanzee calculus from the Taï Forest, Côte d'Ivoire, and human calculus from the Chalcolithic site of Camino del Molino, Spain, we present the first reported observations on characteristics of the matrix setting that are conducive to the survival of starch in dental calculus. We also assess the potential for SEM-EDX to detect starch and differentiate it from structurally and molecularly similar substrates. We demonstrate that SEM-EDX may offer a non-destructive technique for studying microremains in certain contexts. Finally, we compare traditional optical analytical techniques (OM) with less invasive electron microscopy. The results indicate that SEM-EDX and OM are both effective for observing microremains in calculus, but differ in their analytical resolution to identify different microremains, and we therefore recommend a sequential use of both techniques.

## **3.1 Introduction**

Dental calculus, or dental plaque calcified by salivary calcium phosphate, was first noticed as a reservoir of dietary information when Armitage (1975) recognised plant remains on the teeth of archaeological ungulates. Dobney and Brothwell (1986, 1988) later demonstrated the value of calculus in the study of human diets. Analysis of plant and animal microremains in archaeological dental calculus is a rapidly growing field in dietary reconstruction (e.g. Boyadjian et al., 2007; Henry et al., 2012; Liu, 2012; Mickleburgh and Pagán-Jiménez, 2012; Warinner et al., 2014). Researchers have reported starch, phytoliths, pollen, diatoms, chrysophycean cysts, sponge spicules, and mineral particles in human calculus up to tens of thousands of years old (e.g. Dobney and Brothwell, 1988; Boyadjian, 2012).

Despite this interest in dental calculus as a source of dietary information, there are still many questions about the mechanisms by which plant microremains, particularly starch grains, are preserved within the calculus. Native starch grains (i.e. starches in their original, unaltered state) are the major focus of many recent and ancient dietary studies. Starch is a foremost nutritional component in many human and non-human primate diets, and it can also survive in the archaeological record over long periods of time due to its semi-crystalline polysaccharide structure (Mercader et al., 2008; Hardy et al., 2009; Henry et al., 2011; Salazar-García et al., 2013; Leonard et al., 2015). The means by which starch embeds and preserves in calculus is still unclear. The mouth is a hostile environment for starch preservation because of the action of salivary digestive enzymes and bacterial metabolic activity (Lukacs and Largaespada, 2006). Calculus forms gradually as bacteria-rich plaque biofilms mineralise from calcium phosphate in the saliva over a period of days to years (Abraham et al., 2005). During this formation and mineralization process, the starch grains are exposed to  $\alpha$ -amylase, which is present in the saliva of humans and several orders of mammals (Butterworth et al., 2011). Amylase quickly digests starch by breaking down the polysaccharide crystalline structure into various simple and complex sugars through hydrolysis (Lukacs and Largaespada, 2006). Theoretically, starch may avoid oral digestion and survive in protected niche areas in calculus, but this has not yet been empirically confirmed.

In addition to the difficulties with starch preservation in the oral cavity, there is also the possibility that the starches that have been recovered from calculus are actually the result of modern contamination. Modern starches are abundant in the air, water, and working surfaces of most facilities, making environmental contamination a strong possibility. Archaeological and field site contexts suffer from sources of contamination such as airborne starch rain, but the greatest risk of contamination comes from excavation and post-excavation handling in the presence of food or due to the use of gloves powdered with corn or other starches (Newsom and Shaw, 1997; Loy and Barton, 2006; Laurence et al., 2011).

Currently, the standard methodology for starch grain recovery from calculus is too destructive to confirm whether observed starch came from the calculus or from contamination. This method involves mechanically or chemically removing calculus from the tooth, grinding or dissolution to break up the sample, and finally examining the particles using optical light microscopy (OM) (Henry and Piperno, 2008). Furthermore, to the untrained eye, several other calculus components, such as cysts, mineral grains, fungal spores, wood cells, and air bubbles may be confused with starch grains when viewed only under OM. Some have proposed confirming starch presence by measuring amylase activity on treated samples (Hardy et al., 2009), but this enzyme destroys the starch in the process. One common and reliable means to detect starch is to apply iodine potassium iodide (IKI) solution, which binds to the amylose molecule, and look for the characteristic blue-black stain. However, this temporarily obscures the starch's diagnostic surface features. Furthermore, it is impractical to apply a staining solution to an intact calculus matrix because objects within the mineralised matrix are protected from moisture. Accordingly, there is a great need for more sophisticated and non-destructive methods to confirm the successful detection of starch grains in dental calculus. Some researchers have suggested the possibility of using scanning electron microscopy (SEM) to study plant microremains in calculus (Dobney and Brothwell, 1986; Reinhard et al., 2001; Tromp, 2012). Despite the success of this method in locating phytoliths (Arensburg, 1996; Lalueza-Fox et al., 1996; Charlier et al., 2010; Kucera et al., 2011; Tromp, 2012; Tao et al., 2015), the detection of starch grains through SEM has not yet been attempted.

In this study, we present SEM coupled with energy-dispersive X-ray spectroscopy (EDX) as a novel means for identifying starch and other microremains in intact human and chimpanzee dental calculus. This system provides us with the ability to identify microremains, including starch grains, by their morphology and elemental composition *in situ* in the calculus, thus ruling out contamination. It also allows us to explore the kinds of environments within the calculus that may permit starch preservation. Furthermore, we examine the potential of EDX to detect starch by comparing the elemental makeup of native starch to those of saliva-hydrolysed starch and other non-starch saccharides to learn whether EDX distinguishes starch from other polymers of similar elemental makeup. This identification allows us to positively show that starch grains survive in calculus. Finally, we compare the results from SEM-EDX to those from OM on the same human calculus samples to determine whether these techniques offer comparable or complementary results. Due to time constraints, we were unable to conduct this portion of the analysis on the chimpanzee samples and instead only used human dental calculus samples.

## **3.2. Materials and methods**

## *3.2.1 Study groups*

The calculus samples were obtained from two groups, modern wild chimpanzees from the Taï Forest in Côte d'Ivoire and humans from the Chalcolithic collective burial of Camino del Molino in Spain [\(Table 4\)](#page-7-0). We chose these two test groups for the following reasons: 1) individuals from both have abundant calculus on their teeth; 2) they represent modern (chimpanzee) and archaeological (Chalcolithic humans) timeframes; and 3) both groups maintained very different dietary strategies and should therefore have different microremain profiles.

The sample of chimpanzee calculus came from the Taï Chimpanzee Osteology Collection curated at the Max Planck Institute for Evolutionary Anthropology (MPI-EVA) in Leipzig, Germany. The behaviour of the wild chimpanzees living in the Taï Forest has been monitored and documented since the commencement of the Taï Chimpanzee Project in 1979 (Boesch and Boesch-Achermann, 2000). Taï Forest data collection complied with the requirements and guidelines of the Ministère de l'Enseignement Supérieure et de la Recherche Scientifique, and adhered to the legal requirements of Côte d'Ivoire. The osteology collection contains 77 chimpanzees. We chose calculus samples from individuals who had comprehensive observational records documenting diet, sex and age. After their death, the remains of these individuals were interred for defleshing, later exhumed, and curated. We collected calculus from molars or canines of six individuals; two females and four males. The Taï Chimpanzees consume native starch from wild nuts and seeds such as the Gabon nut (*Coula edulis* Baill.) and Kola nut (*Cola nitida* (Vent) Schott et Endl.) (Hohmann et al., 2010; N'guessan, 2012), and unlike humans, they consume no cooked or processed foods. Our preliminary reference collection of Taï Forest chimpanzee foods shows that ten of the 82 foods we analysed are starch-rich. However, these 82 species represent less than a third of plants this population is known to consume, and we are still building this reference collection. Chimpanzees also produce salivary amylase, though purported at much lower quantities than do humans (Perry et al., 2007; Behringer et al., 2013).

Camino del Molino is a Chalcolithic collective burial pit found during construction work in the city of Caravaca de la Cruz (Murcia, southeast Spain). Radiocarbon dates from bone collagen samples spanning the burial sequence indicate that the site was in continual use over a span of 300- 400 years during the first half of the third millennium B.C. The site contained a minimum of 1,300 individuals, likely the remains of 16-20 generations of one population buried at one place (Lomba Maurandi et al., 2009). Approximately 30 % of the individuals are classified as juvenile (<14 yrs.), and the rest are adults spanning from young to old (Haber Uriarte et al., 2013). We collected dental calculus preferably from lower molars for standardisation from the teeth of six individuals; two females, two males, and two individual of unknown sex [\(Table 4\)](#page-7-0). There are no archaeobotanical studies from Camino del Molino or from the broader region of Murcia contemporary to the site. However, studies of Late Neolithic and Chalcolithic deposits in neighbouring regions suggest that the number of cultivated species is low and consists mainly of naked wheat (*Triticum* sp.), barley (*Hordeum vulgare* L.), some lentil (*Lens culinaris*  Medikus*)* and common vetch (*Vicia sativa* L.) (Pérez Jordà, 2005; Pérez Jordà and Carrión Marco, 2011). There is no published study from the site on culinary practices, in part because it is a necropolis and not a habitation site. Despite this, its Chalcolithic age indicates that this population consumed cooked food, because cooking is widespread across the European Neolithic and Bronze Age societies (Thissen et al., 2010; Halstead, 2012).

| Lab identifier | Individual number | Type              | Tooth                   | <b>Sex</b>     | Age (years) | Weight            |
|----------------|-------------------|-------------------|-------------------------|----------------|-------------|-------------------|
| $SI-13-32$     | Sujeto 6          | Camino del Molino | P1 mandible             | F              | $26 - 28$   | $1.76 \text{ mg}$ |
| $SI-13-33$     | Sujeto 8          | Camino del Molino | C maxilla               | М              | $22 - 24$   | $0.51$ mg         |
| $SI-13-36$     | Sujeto 11         | Camino del Molino | I2 mandible             | ?              |             | $6.06$ mg         |
| $SI-13-37$     | Sujeto 17         | Camino del Molino | M <sub>2</sub> mandible | M <sup>2</sup> | 43-55       | $10.0 \text{ mg}$ |
| $SI-13-38$     | Sujeto 113        | Camino del Molino | M <sub>2</sub> mandible | F              | 24-28       | $1.88 \text{ mg}$ |
| $SI-13-39$     | Sujeto 151        | Camino del Molino | C mandible              | M              | 30-35       | $1.09$ mg         |
| Venus          | 15001             | Taï Chimpanzee    | M3 maxilla              | F              | 27          | $0.72 \text{ mg}$ |
| Leo            | 15012             | Taï Chimpanzee    | M <sub>3</sub> mandible | M              | 19          | $1.19 \text{ mg}$ |
| Fanny          | 11780             | Taï Chimpanzee    | M <sub>3</sub> mandible | F              | 25          | $3.34 \text{ mg}$ |
| Goma           | 15004             | Taï Chimpanzee    | M <sub>3</sub> mandible | F              | 28          | $2.40$ mg         |
| Rubra          | 15023             | Taï Chimpanzee    | M <sub>3</sub> mandible | F              | 38?         | $3.88$ mg         |
| Castor         | 13439             | Taï Chimpanzee    | M <sub>3</sub> mandible | F              | 22          | $2.25$ mg         |

<span id="page-7-0"></span>**Table 4: Calculus samples analysed using SEM-EDX and OM. Sex and age classification of the Camino del Molino remains are preliminary (Haber Uriarte et al., 2013).**

## *3.2.2 Calculus sampling*

We selected teeth encrusted with a prominent band of calculus present on the enamel surface. We sampled only supragingival calculus (above the gum line), since it is unclear if subgingival calculus (below the gum line, on the neck of the tooth) preserves food remains. We photographed the calculus before sampling, and then brushed the sample tooth gently with a dry, sterile toothbrush to remove surface contaminants. We then used a dental scalar to remove small areas of supragingival calculus (~4 mm area), from the enamel. We conducted all calculus sampling in a positive pressure hood at the archaeological science laboratories at the MPI-EVA. We then weighed each of the samples and transferred them to microcentrifuge tubes for storage until further use. Following sampling, the teeth and surviving calculus were photographed again. Additionally, we collected control samples, including the packing material in which the teeth had been stored.

#### *3.2.3 Electron microscopy analysis*

We conducted the SEM-EDX analysis at University College Dublin's Nano-Imaging and Materials Analysis Centre (NIMAC) in Dublin, Ireland. The calculus samples were mounted on stubs using double-sided carbon tape, and sputter coated with gold for 20 seconds using an Emitech K575X Sputter Coating Unit, to prevent surface charging by the electron beam. We then examined the calculus using a FEI Quanta 3D FEG DualBeam (FEI Ltd, Hillsboro, USA) SEM with an attached EDAX ED APOLLO XV Silicon Drift Detector with a 5 – 10 kV accelerating voltage. EDX detected and documented most elements of interest excluding hydrogen, which is non-detectable with this method. We omitted the gold elemental peak from each spectrum since the gold was added during sputter coating. We photographed and documented every tentative microremain and later described our observations.

## *3.2.4 Carbohydrate reference standards and partially hydrolysed controls*

We used a variety of reference standards (see [Table 5\)](#page-8-0) to assess the accuracy of EDX reads on the experimental sample types of starch. Starches from a variety of plants were selected to represent major starch types such as corn starch, potato starch, and common dietary components for each population (Boesch and Boesch 1983): wheat (*Triticum aestivum* L.), Gabon nut (*Coula edulis* [Baill.\)](http://en.wikipedia.org/wiki/Henri_Ernest_Baillon), Xylia (*Xylia evansii* Hutch.) and Kola nut (*Cola nitida* [Vent] Schott et Endl.). The nuts were ground, dried and weighed to derive nut flour suitable for use. Wheat, potato and corn were purchased from local distributers in Germany (see [Table 5\)](#page-8-0).

<span id="page-8-0"></span>**Table 5: List of reference samples analysed using EDX.**



Laboratory-grade fructose, maltose, glucose and sucrose (Roth, Germany) were included as standards because they have nearly identical elemental compositions as starch but with structurally different molecular arrangements (e.g. sucrose has 2.1 wt % (mass fraction) more carbon than fructose, but 2.1 wt % less than starch).

To compare EDX element signatures for the different types of saccharides, we took EDX measurements from five individual grains of fructose, sucrose, maltose, glucose and wheat starch, corn starch, Kola nut starch, Xylia starch and potato starch. This allows the comparison of a monosaccharide, a disaccharide, and a polysaccharide (starch).

Finally, to assess whether EDX signatures and detection accuracy is affected by the salivary modification (hydrolysis) of starch, we experimentally hydrolysed the native starches from the wheat flour and both nut varieties using salivary amylase derived from human saliva – a simulation of the effects of oral digestion on starch that can occur. One of us (R.C.P.) provided the saliva used in all experiments, which was collected on a single occasion. We split each of the individual plant samples into nine subsamples of approximately 2 mg each: three subsample per plant remained untreated (control), three were exposed to amylase  $(35 \mu L)$  of saliva) for 30 minutes, and three were exposed to amylase  $(35 \mu L)$  of saliva) for 90 minutes. We also similarly partitioned the wheat flour into nine aliquots into three subsamples of 2 mg each for identical amylase treatment. We ceased hydrolysis by displacing the saliva with alcohol and centrifugation at 1691 x g to remove as much liquid from the sample as possible and stop hydrolysis. Then the remaining alcohol was evaporated at  $35\text{ }^{\circ}$  C in a drying oven. We performed measurements using SEM-EDX in triplicate on one starch grain from each subsample, creating nine readings per category (e.g. wheat 30 minute hydrolysed). A summary of these analyses is provided in Figs. 3 and 4.

## *3.2.5 Optical microscopy analysis*

We performed optical microscopy on the ancient human remains at the Plants Working Group Laboratory in the MPI-EVA, Leipzig. We removed the gold plated calculus samples from the SEM mounting stubs, and then ground them in a 1.5 ml Eppendorf microcentrifuge tube with a micro pestle containing  $\sim$  50  $\mu$ l of a 25 % glycerine solution to reduce sample loss due to static electricity. Glycerine was chosen as its refractive index is lower than starch making it suitable for starch detection. The samples were then centrifuged at 1691 x g (Heraeus MEGAFUGE 16 with TX-400 Swinging Bucket Rotors) for 10 minutes. All of the resulting pellets were mounted on glass slides and examined under bright field and cross-polarised light on an A1 Zeiss Axioscope microscope at 400 × magnification. Larger samples were mounted on several slides. Each microremain was photographed and described (Table 6).

#### **3.3. Results**

#### *3.3.1 Standards*

The EDX spectrum of starch is distinct from other saccharides but not sufficiently to permit reliable identification [\(Fig.](#page-11-0) 3, [Fig.](#page-11-1) 4; Appendix table 1, Appendix table 2). The EDX results from all the samples indicate that oxygen is underrepresented. Though carbon comprises roughly 40-50 wt % of these saccharides, the EDX spectra indicates carbon at 60-90 wt % [\(Fig.](#page-11-0) 3). Comparing the short-chain saccharides to the starches there is a difference but some types of starch overlaps with each short-chain saccharides. This indicates that some starch may be distinguishable from short-chain saccharides through EDX. There was far more variability in carbon values in starch than in short-chain saccharides. Starch is composed of oxygen, hydrogen and carbon  $(C<sub>6</sub>H<sub>10</sub>O<sub>5</sub>)<sub>n</sub>$ , where n ranges from 300 to 1000, so starch is approximately 42.1 %=carbon, 6.5 %=hydrogen, 51.4 %=oxygen (Newman et al., 1996). Thus, maize starch comes closest to the expected values of starch. This variability possibly reflects the heterogeneous nature of the starch. Starch varies in both proportion of amylose and amylopectin and minor compounds such as proteins and lipids (Belitz et al., 2009). We see further evidence of this elemental variability in the native starch samples in [Fig.](#page-11-1) 4, which had a higher variability of both carbon and oxygen than the hydrolysed starches. The EDX profiles of hydrolysed starches fall within the range of their native counterparts, yet they show noticeable less variation and reduced oxygen values [\(Fig.](#page-11-1) 4). The reduction in variation and lower oxygen levels in these samples may be either from the result of the added ethanol reducing oxygen in the starch or the ethanol washed off debris on the starch surface. A few of the damaged starches have slightly increased oxygen percentages, but this is not consistent across all hydrolysed subsamples. We found no evidence that saliva-activated hydrolysis could obscure the starch EDX elemental signature. However, when large starch shaped objects are present under SEM, it is possible to test whether these particles have a molecular make-up that is similar to starch and other saccharides.



<span id="page-11-0"></span>**Fig. 3: Carbon wt % (mass fraction) of starch, sugars produced by hydrolysis and reference sugars. Plot shows five individual grains of starches, glucose, and maltose and out-group carbohydrates (fructose and sucrose) detected with EDX. Values exclude contaminating elements such as potassium from sweat (3.6.1).**



<span id="page-11-1"></span>**Fig. 4:** C**omparing native starch versus samples that were hydrolysed with amylase for 30 and 90 minutes at room temperature. Three starches were sampled with triplicate readings. Values exclude contaminating elements such as potassium such as potassium from sweat (3.6.2).**

#### *3.3.2 Calculus samples*

Examination of the SEM images of the calculus confirmed that this matrix has a heterogeneous texture, with smooth surfaces as well as many pores, crack and crevices [\(Fig.](#page-13-0) 5, [Fig.](#page-14-0) 6). Most of the pores appear to be the result of rod bacterial pseudomorphs, which are shallow and measure only between 0.3 -1 µm in width (bottom left in [Fig.](#page-13-0) 5, and widely scattered in [Fig.](#page-14-0) 6). These pores are too small to preserve microremains, but larger cracks and crevices had many microremains [\(Fig.](#page-14-0) [6\)](#page-14-0). Further examination of the calculus revealed several types of inclusions within the matrix. In some cases, these inclusions were consistent with the overall size (15- 40 μm) and shape (ovoid- pyriform) of certain starch grain types, and inconsistent with other microremains such as yeast and bacterial cells [\(Fig.](#page-13-0) 5). The supposed starch clusters were clearly embedded in the matrix, with grains occluded by overlying deposits of the matrix material. Interestingly, the starch grains were not evenly distributed in the matrix, and often appeared in clumps [\(Fig.](#page-13-0) 5). This could be explained in two ways; i) plant microremains are deposited in groups originating from clumps in food lumps, or ii) microremains are only preserved in localised niches, such as larger cracks and crevices, in the calculus matrix.



<span id="page-13-0"></span>**Fig. 5: Starch grains located** *in-situ* **on dental calculus surface. SEM image showing a group of starches trapped in the matrix of one of the chimpanzee dental calculus samples (Venus), with the corresponding EDX spectrum (right) showing a calcium phosphate and silicon mantle covering a carbon rich starch (A) and solely a carbon-rich starch (B).**



<span id="page-14-0"></span>**Fig. 6: SEM image showing microremain diversity. A concentration of pollen (A) and sponge spicules (B) in SJ-13-39 from Camino del Molino. Microremains were often found clustered.**



<span id="page-14-1"></span>**Fig. 7: SEM image showing localised damage that arises from higher primary voltage SEM (10 kV) and EDX on a spicule in calculus from Camino del Molino. Before (left); after (right).**

The EDX spectra of the calculus matrix from all of our samples indicate that it is mostly composed of calcium and phosphorus, with trace amounts of aluminium, magnesium, silicon, sodium and manganese (Appendix table 3). These elements confirm our supposition that the majority of our samples consist of calculus, a mixture of hydroxyapatite and other minerals, rather than contaminating exogenous matter (Charlier et al., 2010; Salazar-García et al., 2014). In some instances, silicon was locally abundant in the calculus (Appendix table 3), which may be important for the preservation of starch grains. In contrast to the mineral matrix, the suspected starch clusters, such as on chimpanzees Venus and Castor had significant carbon peaks. Additionally, the starches often had calcium and phosphorus peaks, reinforcing visual observations that they were indeed embedded in calculus [\(Fig.](#page-13-0) 5). The combination of shape and elemental data [\(Fig.](#page-13-0) 5) is strongly suggestive of *in-situ* findings of microremains preserved in the dental calculus environment. This is possible as starch is morphologically distinct from other carbon rich particles such as fungal filaments, *Candida albicans* cells, cellulose and sugars. We also note that the starch we located with SEM-EDX was undamaged and we did not locate any semi gelatinised or hydrolysed starch. These were also not observed with optical microscopy but it is possible damaged starch was not visible with this approach.

In addition to the starches, we also identified a variety of other plant and animal microremains preserved in the calculus using SEM-EDX, including phytoliths, sponge spicules, diatoms and pollen (Table 6). These microremains were identified by their diagnostic morphology using conventional methods (Torrence and Barton, 2006; Nadel et al., 2013; Power et al., 2014a), and this identification was confirmed by their EDX spectra. For example, spicules were easily identified based on their long rectangular shape and high level of regularity [\(Fig.](#page-14-0) 6 and [Fig.](#page-14-1) 7) unlike smooth longcell phytoliths, and EDX readings confirmed their biogenic silica composition (7.1). OM also demonstrated the presence of a rich assemblage of plant microremains [\(Table 5\)](#page-8-0). We noted some of these microremains during the SEM analysis, such as the abundant monaxon spicules [\(Fig.](#page-14-0) 6), but we only detected some, such as multicell long-cell phytoliths, unsilicified plant cells and calcium oxalate [\(Fig.](#page-17-0) 8), with OM.



## **Table 6: Recovered microremains using both microscopy approaches.**

+ : a high but unquantified number



**Fig. 8: A calcium oxalate prism observed with optical microscopy in SJ-13-37; under bright field optical microscopy (left), and cross-polarising optical microscopy (right).**

<span id="page-17-0"></span>A comparison of the microremains observed under SEM-EDX with those seen in OM revealed important differences (Table 6). We observed more starch microremains using OM than SEM-EDX. This is probably because the sample preparation for OM breaks down the calculus matrix, freeing starch microremains that were trapped in the middle of the calculus chunk. Yet paradoxically, other microremains, such as sponge spicules, were more commonly seen in SEM-EDX than in OM of the same samples.

Based solely on the SEM results (we did not perform OM on the chimpanzee samples), the two groups we studied did present some differences. The chimpanzee samples were rich in starch grains and diatoms, while the human samples had an abundance of unsilicified plant cells and sponge spicules (Table 6).

#### **3.4. Discussion**

Analysis of calculus samples by SEM-EDX and OM provides data that validates the study of microremains recovered from this biological material. By SEM-EDX, we were able to identify the elemental constituents of starch, and confirm its position *in situ* in calculus particles. This is the first time that starch has been identified by its elemental signature while still embedded within the calculus matrix, and confirms that starch can be preserved in calculus, and can therefore be a useful source of dietary information.

The analysis suggests that certain features of the calculus may promote the preservation of microremains, and starch grains in particular. Food debris may trap around calculus rather than calculus growing around food debris. While the pores left from bacteria colonies were too small to provide a protected niche for starches, larger cracks and crevices were full of microremains, possibly because these areas provided a protected environment. Furthermore, the silicon we detected in the dental calculus may be significant. Silicic acid can induce spontaneous precipitation of calcium phosphate in the saliva, which is the precursor mineral necessary for calculus formation. Silicic acid may be consumed directly via water or indirectly via plants, as it enters plants along with groundwater. Consuming polysilicic acid and silica increases calculus formation, thereby regulating this process (Damen and Ten Cate, 1989; Roberts-Harry and Clerehugh, 2000; Jin and Yip, 2002). Our observations of silicon concentrations adjacent to embedded starch clusters [\(Fig.](#page-13-0) 5) corroborates these reports, suggesting that dietary exposure to silica or silicic acid enables enhanced calculus formation and thus the preservation of native starch in dental calculus.

By following the SEM analysis with an OM examination of the same samples, we are able to compare the effectiveness of each for specific microremain types. Sponge spicules were easily visualised under SEM, but were seen less with OM. This may be because the spicules are relatively fragile and are damaged when the calculus is processed, possibly explaining why spicules are rarely reported in dental calculus studies (Tromp, 2012; Dudgeon and Tromp, 2014). Because these particles, as well as diatoms and Chrysophycean cysts, are highly dependent upon water sources, they may indicate source type and provenance of consumed water, making them powerful potential ecological markers for primatology and archaeology studies (Dudgeon and Tromp, 2014). In contrast, calcium oxalate crystals were only visible under OM, and not SEM. These crystals, which may occur as druses, raphides or other similar forms, are a potentially useful marker of plants. They may be more visible using OM because they have high interference colours that are visible under cross polarised light [\(Fig.](#page-17-0) 8). For reasons that remain unclear, calcium oxalate is rarely reported or discussed in calculus literature. Some research indicates that calcium oxalate does not survive due to acidity in the mouth (Tromp, 2012), but given their sheer abundance in plants and the relatively neutral oral pH, it is likely that calcium oxalates do survive and are simply overlooked. On the other hand,

starch grains were clearly visible using both SEM and OM. However, we did note that within individuals, the starches that we observed under OM typically did not match the size and morphology of those seen in SEM-EDX. This contrasts with the spicules, which often matched size and shape. This is likely due to the small number of starches but high number of spicules. We also observed pollen grains embedded in the calculus using SEM [\(Fig.](#page-14-0) 6) and OM. Although this type of pollen grain were too small to analyse with the EDX, we do believe that SEM-EDX may be appropriate for identifying many larger types of pollen grains, since these plant remains are composed of potassium, magnesium, sodium and calcium (Szczęsna, 2007) and should be easily visible in the EDX spectra.

Finally, the SEM analysis accurately reflected some stark differences between our study groups. The differences in microremains number and types between the chimpanzee and humans likely reflect the dietary behaviour and the age of the remains. The chimpanzees consumed only raw plants, while the human group potentially cooked much of their food. The chimpanzees therefore consumed many more native, undamaged starch grains, and so there is greater opportunity for the preservation of native starch grains in dental calculus. Though the humans may have consumed more starch overall, many of these starches would have been semi gelatinised through cooking, disrupting the semi-crystalline structure and reducing the potential for starch preservation in the mouth (Holm et al., 1988). Cooking, combined with higher levels of salivary amylase in humans relative to chimpanzees (Perry et al., 2007; Behringer et al., 2013) may have greatly reduced the relative proportion of starch entering the human calculus matrix during its formation. Furthermore, the chimpanzee samples are modern and likely to be well-preserved while starch in the human calculus may have depleted due to digenesis over thousands of years.

Overall, SEM-EDX does allow us to visualise and identify microremains embedded in dental calculus, but this technique is not without limitations and constraints. Internal features of starch grains that are vital for identifying the taxonomic origin of the starch are not visible under SEM. We found that when using EDX combined with higher primary voltage (10 kV), the beam moved or damaged fragile microremains such as spicules [\(Fig.](#page-14-1) 7). EDX can only give reliable data on objects ≥4 μm due to the penetration of the beam, making it impossible to measure very small microremains including smaller starches. We found other techniques such as backscatter detection to be of little additional advantage in detecting starch, though this method may be useful in certain contexts such as examining calculus for

embedded phytoliths (Tromp, 2012). It is possible to examine only the surface portion of intact calculus matrix using SEM-EDX, and so this is not a viable method for visualising interior dental calculus structure and microremains. Sample preparation may also be destructive since samples must be gold-plated and mounted, but use of SEM without the plating may cause the sample image quality and identification power to deteriorate.

## **3.5. Conclusions**

The visual identification and subsequent elemental testing of microremains embedded in the dental calculus of humans and chimpanzees suggests that these important dietary markers are indeed trapped and preserved in calculus during the lifetime of the individual. Clearly, this matrix has a protective quality that shields fragile and degradable components, namely starch, from the enzymatic oral environment.

SEM-EDX and OM have different sensitivity to different microremains. SEM-EDX offers a means to confirm the presence of starch by combining morphological and elemental information without having to destroy either the calculus, as required in processing for OM, or the starch grains themselves, as proposed when using enzymatic reactions. Even if starch is semi gelatinised it should preserve an elemental signature that is suggestive of starch. We applied SEM-EDX to intact calculus to witness microremains *in situ*, but this technique is equally viable for more finely processed calculus samples mounted on plates, or even to calculus still attached on the original tooth. However, it is important to note that diagnostic features of starch grains, such as the hilum and lamellae, are only visible using OM.

Our study indicates that SEM-EDX is a viable alternative to OM analysis of calculus, but researchers should choose their analytical method based on the questions they seek to answer, and the plant microremains that they intend to study. Furthermore, on very sensitive osteological remains, it may be possible to use SEM-EDX to study calculus using entirely non-destructive means to examine embedded microremains directly on the tooth; a useful technique if the tooth is not firmly attached in the mandible or maxilla. We prefer to consider SEM-EDX a complementary rather than replacement technique in the study of dental calculus microremains. A sequential workflow that first examines calculus under SEM-EDX and then under OM may be the optimal solution for highest resolution of microremains, though we recognise that this approach is time intensive and can be

costly. We believe that further exploration and experimentation of SEM techniques is important in the field of archaeological and palaeodietary reconstruction. The continued refinement and expansion of dental calculus analysis techniques is an important focus in order to optimise the information we can harvest from this ephemeral and fragile material.